SEQUENCE LISTING

(1) GENERAL INFORMATION:

- (i) APPLICANT: BYWATER, MARGARET LINDSTROM, PER INGANAS, MATS
- (ii) TITLE OF INVENTION: SEQUENCE-BASED MUTATION ANALYSIS OF NEOPLASTIC TISSUE FOR DIAGNOSIS OR PROGNOSIS OF THE NEOPLASIA
- (iii) NUMBER OF SEQUENCES: 25
- (iv) CORRESPONDENCE ADDRESS:
 - (A) ADDRESSEE: BIRCH, STEWART, KOLASCH AND BIRCH
 - (B) STREET: PO BOX 747
 - (C) CITY: FALLS CHURCH
 - (D) STATE: VA
 - (E) COUNTRY: USA
 - (F) ZIP: 22040-0747
- (v) COMPUTER READABLE FORM:
 - (A) MEDIUM TYPE: Floppy disk
 - (B) COMPUTER: IBM PC compatible
 - (C) OPERATING SYSTEM: PC-DOS/MS-DOS
 - (D) SOFTWARE: PatentIn Release #1.0, Version #1.30

(vi) CURRENT APPLICATION DATA:

- (A) APPLICATION NUMBER:
- (B) FILING DATE:
- (C) CLASSIFICATION:

(viii) ATTORNEY/AGENT INFORMATION:

- (A) NAME: MURPHY JR, GERALD M
- (B) REGISTRATION NUMBER: 28,977
- (C) REFERENCE/DOCKET NUMBER: 1614-178

(ix) TELECOMMUNICATION INFORMATION:

- (A) TELEPHONE: (703) 205-8000
- (B) TELEFAX: (703) 205-8050

(2) INFORMATION FOR SEQ ID NO:1:

- (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 19 base pairs
 - (B) TYPE: nucleic acid
 - (C) STRANDEDNESS: single
 - (D) TOPOLOGY: linear
- (ii) MOLECULE TYPE: other nucleic acid
 - (A) DESCRIPTION: /desc = "PRIMER"

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	(xi)	SEQUENCE DESCRIPTION: SEQ ID NO:1:	
GT	GCTTTC	CCA CGACGGTG A	19
(2)) INFO	DRMATION FOR SEQ ID NO:2:	
	(i)	SEQUENCE CHARACTERISTICS: (A) LENGTH: 23 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: single (D) TOPOLOGY: linear	
	(ii)	MOLECULE TYPE: other nucleic acid (A) DESCRIPTION: /desc = "PRIMER"	
	(xi)	SEQUENCE DESCRIPTION: SEQ ID NO:2:	
TC	ATGTGC	CTG TGACTGCTTG TAG	23
(2)) INFO	DRMATION FOR SEQ ID NO:3:	
((i)	SEQUENCE CHARACTERISTICS: (A) LENGTH: 19 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: single (D) TOPOLOGY: linear	
	(ii)	MOLECULE TYPE: other nucleic acid (A) DESCRIPTION: /desc = "PRIMER"	
	(xi)	SEQUENCE DESCRIPTION: SEQ ID NO:3:	
GG	CGGGGG	GTG TGGAATCAA	19
(2) INFO	DRMATION FOR SEQ ID NO:4:	
	(i)	SEQUENCE CHARACTERISTICS: (A) LENGTH: 23 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: single (D) TOPOLOGY: linear	
	(ii)	MOLECULE TYPE: other nucleic acid (A) DESCRIPTION: /desc = "PRIMER"	

(xi) SEQUENCE DESCRIPTION: SEQ ID NO:4:

TCTGGCATTC TGGGAGCTTC ATC 23		
(2) INFORMATION FOR SEQ ID NO:5:		
 (i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 20 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: single (D) TOPOLOGY: linear 		
<pre>(ii) MOLECULE TYPE: other nucleic acid (A) DESCRIPTION: /desc = "PRIMER"</pre>		
(xi) SEQUENCE DESCRIPTION: SEQ ID NO:5:		
CCGTCCCAGT AGATTACCAC	20	
(2) INFORMATION FOR SEQ ID NO:6:		
 (i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 20 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: single (D) TOPOLOGY: linear 		
<pre>(ii) MOLECULE TYPE: other nucleic acid (A) DESCRIPTION: /desc = "PRIMER"</pre>		
(xi) SEQUENCE DESCRIPTION: SEQ ID NO:6:		
GTTTTCCGTC TGGGCTTCTT	20	
(2) INFORMATION FOR SEQ ID NO:7:		
 (i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 27 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: single (D) TOPOLOGY: linear 		
<pre>(ii) MOLECULE TYPE: other nucleic acid (A) DESCRIPTION: /desc = "PRIMER"</pre>		
(xi) SEQUENCE DESCRIPTION: SEQ ID NO:7:		
CTGTGACTTG CACGTACTCC CCTGCCC	27	

(2) INFORMATION FOR SEQ ID NO:8:

	i)	i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 23 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: single (D) TOPOLOGY: linear	
	(ii	i) MOLECULE TYPE: other nucleic acid (A) DESCRIPTION: /desc = "PRIMER"	
	(xi	i) SEQUENCE DESCRIPTION: SEQ ID NO:8:	
	TAGACTO	SACC CTTTTTGGAC TTC	23
	(2) INE	FORMATION FOR SEQ ID NO:9:	
((i	i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 21 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: single (D) TOPOLOGY: linear	
	(ii	i) MOLECULE TYPE: other nucleic acid (A) DESCRIPTION: /desc = "PRIMER"	
	(xi	i) SEQUENCE DESCRIPTION: SEQ ID NO:9:	
	CGTGTG	GAGT ATTTGGATGA C	21
	(2) INE	FORMATION FOR SEQ ID NO:10:	
	i)	i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 22 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: single (D) TOPOLOGY: linear	
	(ii	i) MOLECULE TYPE: other nucleic acid (A) DESCRIPTION: /desc = "PRIMER"	
	(xi	i) SEQUENCE DESCRIPTION: SEQ ID NO:10:	
	TGGTTTC	CTTC TTTGGCTGGG GA	22
	(2) INE	FORMATION FOR SEQ ID NO:11:	
	(i	i) SEQUENCE CHARACTERISTICS:	

(A) LENGTH: 21 base pairs

	(B) TYPE: nucleic acid(C) STRANDEDNESS: single(D) TOPOLOGY: linear	
	<pre>(ii) MOLECULE TYPE: other nucleic acid (A) DESCRIPTION: /desc = "PRIMER"</pre>	
	(xi) SEQUENCE DESCRIPTION: SEQ ID NO:11:	
GCT	TTGAGGT GCGTGTTTGT G	21
(2)	INFORMATION FOR SEQ ID NO:12:	
	 (i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 21 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: single (D) TOPOLOGY: linear 	
	<pre>(ii) MOLECULE TYPE: other nucleic acid (A) DESCRIPTION: /desc = "PRIMER"</pre>	
	(xi) SEQUENCE DESCRIPTION: SEQ ID NO:12:	
CTG'	TCAGTGG GGAACAAGAA G	21
(2)	INFORMATION FOR SEQ ID NO:13:	
	 (i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 20 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: single (D) TOPOLOGY: linear 	
	<pre>(ii) MOLECULE TYPE: other nucleic acid (A) DESCRIPTION: /desc = "PRIMER"</pre>	
	(xi) SEQUENCE DESCRIPTION: SEQ ID NO:13:	
CCN/	GCACTAA GCGAGCACTG	20
		20
(2)	INFORMATION FOR SEQ ID NO:14:	
	 (i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 21 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: single (D) TOPOLOGY: linear 	

	<pre>(ii) MOLECULE TYPE: other nucleic acid (A) DESCRIPTION: /desc = "PRIMER"</pre>	
	(xi) SEQUENCE DESCRIPTION: SEQ ID NO:14:	
CAC	ACGCTTC CCTGGATTGG C	21
		21
(2)	INFORMATION FOR SEQ ID NO:15:	
	(i) SEQUENCE CHARACTERISTICS:(A) LENGTH: 22 base pairs(B) TYPE: nucleic acid(C) STRANDEDNESS: single(D) TOPOLOGY: linear	
121	<pre>(ii) MOLECULE TYPE: other nucleic acid (A) DESCRIPTION: /desc = "PRIMER"</pre>	
P	(xi) SEQUENCE DESCRIPTION: SEQ ID NO:15:	
GCA)	AAACATC TTGTTGAGGG CA	22
(2)	INFORMATION FOR SEQ ID NO:16:	
	 (i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 24 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: single (D) TOPOLOGY: linear 	
	<pre>(ii) MOLECULE TYPE: other nucleic acid (A) DESCRIPTION: /desc = "PRIMER"</pre>	
	(xi) SEQUENCE DESCRIPTION: SEQ ID NO:16:	
CAG	GGGAGTA CGTGCAAGTC ACAG	24
(2)	INFORMATION FOR SEQ ID NO:17:	
	 (i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 22 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: single (D) TOPOLOGY: linear 	
	<pre>(ii) MOLECULE TYPE: other nucleic acid (A) DESCRIPTION: /desc = "PRIMER"</pre>	

	((xi) SEQUENCE DESCRIPTION: SEQ ID NO:17:	
	GTTTC	CCGTCT GGGCTTCTTG CA	22
	(2) I	INFORMATION FOR SEQ ID NO:18:	
		 (i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 21 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: single (D) TOPOLOGY: linear 	
	(<pre>(ii) MOLECULE TYPE: other nucleic acid (A) DESCRIPTION: /desc = "PRIMER"</pre>	
1			
	((xi) SEQUENCE DESCRIPTION: SEQ ID NO:18:	
	GGTAC	CAGTCA GAGCCAACCT C	21
	(2) I	INFORMATION FOR SEQ ID NO:19:	
		 (i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 21 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: single (D) TOPOLOGY: linear 	
	((ii) MOLECULE TYPE: other nucleic acid (A) DESCRIPTION: /desc = "PRIMER"	
	((xi) SEQUENCE DESCRIPTION: SEQ ID NO:19:	
	GCCAA	ACCTCA GGCGGCTCAT A	21
	(2) I	INFORMATION FOR SEQ ID NO:20:	
		(i) SEQUENCE CHARACTERISTICS:(A) LENGTH: 21 base pairs(B) TYPE: nucleic acid(C) STRANDEDNESS: single(D) TOPOLOGY: linear	
	(<pre>(ii) MOLECULE TYPE: other nucleic acid (A) DESCRIPTION: /desc = "PRIMER"</pre>	

(xi) SEQUENCE DESCRIPTION: SEQ ID NO:20:	
TGGCC	CCCTCC TCAGCATCTT A	21
(2) I	NFORMATION FOR SEQ ID NO:21:	
	 (i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 20 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: single (D) TOPOLOGY: linear 	
((A) DESCRIPTION: /desc = "PRIMER"	
((xi) SEQUENCE DESCRIPTION: SEQ ID NO:21:	
CAAGG	GCCTCA TTCAGCTCTC	20
(2) I	NFORMATION FOR SEQ ID NO:22:	
	(i) SEQUENCE CHARACTERISTICS:(A) LENGTH: 21 base pairs(B) TYPE: nucleic acid(C) STRANDEDNESS: single(D) TOPOLOGY: linear	
. ((ii) MOLECULE TYPE: other nucleic acid (A) DESCRIPTION: /desc = "PRIMER"	
((xi) SEQUENCE DESCRIPTION: SEQ ID NO:22: '	
CGAGT	GGAAG GAAATTTGCG T	21
(2) I	NFORMATION FOR SEQ ID NO:23:	
	 (i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 22 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: single (D) TOPOLOGY: linear 	
((ii) MOLECULE TYPE: other nucleic acid (A) DESCRIPTION: /desc = "PRIMER"	

(xi) SEQUENCE DESCRIPTION: SEQ ID NO:23:

22 CGGCGCACAG AGGAAGAGAA TC (2) INFORMATION FOR SEQ ID NO:24: (i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 23 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: single (D) TOPOLOGY: linear (ii) MOLECULE TYPE: other nucleic acid (A) DESCRIPTION: /desc = "PRIMER" (xi) SEQUENCE DESCRIPTION: SEQ ID NO:24: CGCACACCTA TTGCAAGCAA GGG 23 (2) INFORMATION FOR SEQ ID NO:25: (i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 21 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: single (D) TOPOLOGY: linear (ii) MOLECULE TYPE: other nucleic acid (A) DESCRIPTION: /desc = "PRIMER" (xi) SEQUENCE DESCRIPTION: SEQ ID NO:25:

21

GGGGAGCCTC ACCACGAGCT G